



FIGURE 1

1 ACCCACGCGC AGCGGCCGGA GATGCAGCGG GCGGCCGCGC TGTGCCTGCG ACTGTGGCTC
61 TGCCTGGGAC TCCTGGACGG CCTGGTGAGT GACTACTCCA TGACCCCCC GACCTTGAAC
121 ATCACGGAGG AGTCACACGT CATCGACACC GGTGACAGCC TGTCCATCTC CTGCAGGGGA
181 CAGCACCCCC TCAGAGTGGGC TTGGCCAGGA GCTCAGGAGG CGCCAGCCAC CGGAGACAAG
241 GACAGCGAGG ACACGGGGGT GGTGCGAGAC TGCAGGGGCA CAGACGCCAG GCCCTACTGC
301 AAGGTGTTGC TGCTGCACGA GGTACATGCC AACGACACAG GCAGCTACGT CTGCTACTAC
361 AAGTACATCA AGGCACGCAT CGAGGGCACC ACGGCCGCCA GCTCCTACGT GTTCGTGAGA
421 GACTTTGAGC AGCCATTTCAT CAACAAGCCT GACACGCTCT TGGTCAACAG GAAGGACGCC
481 ATGTGGGTGC CCTGTCTGGT GTCCATCCCC GGCCTCAATG TCACGCTGCG CTCGCAAAGC
541 TCGGTGCTGT GGCCAGACGG GCAGGAGGTG GTGTGGGATG ACCGGCGGGG CATGCTCGTG
601 TCCACGCCAC TGCTGCACGA TGCCCTGTAC CTGCAGTGCG AGACCACCTG GGGAGACCAG
661 GACTTCCTTT CCAACCCCTT CCTGGTGAC ATCACAGGCA ACGAGCTCTA TGACATCCAG
721 CTGTTGCCCA GGAAGTCGCT GGAGCTGCTG GTAGGGGAGA AGCTGGTCCT CAACTGCACC
781 GTGTGGGCTG AGTTTAACTC AGGTGTCACC TTTGACTGGG ACTACCCAGG GAAGCAGGCA
841 GAGCGGGGTA AGTGGGTGCC CGAGCGACGC TCCCAACAGA CCCACACAGA ACTCTCCAGC
901 ATCCTGACCA TCCACAACGT CAGCCAGCAC GACCTGGGCT CGTATGTGTG CAAGGCCAAC
961 AACGGCATCC AGCGATTTTCG GGAGAGCACC GAGGTCATTG TGCATGAAAA TCCCTTCATC
1021 AGCGTCGAGT GGCTCAAAGG ACCCATCCTG GAGGCCACGG CAGGAGACGA GCTGGTGAAG
1081 CTGCCCCTGA AGCTGGCAGC GTACCCCCCG CCCGAGTTCC AGTGGTACAA GGATGGAAAG
1141 GCACTGTCCG GCGGCCACAG TCCCATGCC CTGGTGCTCA AGGAGGTGAC AGAGGCCAGC
1201 ACAGGCACCT ACACCCCTCG CCTGTGGAAC TCCGCTGCTG GCCTGAGGCG CAACATCAGC
1261 CTGGAGCTGG TGGTGAATGT GCGCCCCAG ATACATGAGA AGGAGGCCTC TCCCCCAGC
1321 ATCTACTCGC GTACAGCCG CCAGGCCCTC ACCTGCACGG CCTACGGGGT GCCCCCTGCCT
1381 CTCAGCATCC AGTGGCAC TGCGGCCCTG ACACCCCTGCA AGATGTTTGC CCAGCGTAGT
1441 CTCCGGCGGC GGCAGCAGCA AGACCTCATG CCACAGTGCC GTGACTGGAG GCGGGTGACC
1501 ACGCAGGATG CCGTGAACCC CATCGAGAGC CTGGACACCT GGACCGAGTT TGTGGAGGGA
1561 AAGAATAAGA CTGTGAGCAA GCTGGTGATC CAGAATGCCA ACGTGTCTGC CATGTACAAG
1621 TGTGTGGTCT CCAACAAGGT GGGCCAGGAT GAGCGGCTCA TCTACTTCTA TGTGACCACC
1681 ATCCCCGACG GCTTCACCAT CGAATCCAAG CCATCCGAGG AGCTACTAGA GGGCCAGCCG
1741 GTGCTCCTGA GCTGCCAAGC CGACAGCTAC AAGTACGAGC ATCTGCGCTG GTACCGCCTC
1801 AACCTGTCCA CGCTGCACGA TGCGCACGGG AACCCGCTTC TGCTCGACTG CAAGAACGTG
1861 CATCTGTTTC CCACCCCTCT GGCCGCCAGC CTGGAGGAGG TGGCACCTGG GCGCGGCCAC
1921 GCCACGCTCA GCCTGAGTAT CCCCCGCGTC GCGCCCGAGC ACGAGGGCCA CTATGTGTGC
1981 GAAGTGCAAG ACCGGCGCAG CCATGACAAG CACTGCCACA AGAAGTACCT TCTGGTGAGC
2041 GCCCTGGAAG CCCCTCGGCT CACGCAGAAC TTGACCGACC TCCTGGTGAA CGTGAGCGAC
2101 TCGCTGGAGA TGCAAGTCTT GGTGGCCGGA GCGCACGCGC CCAGCATCGT GTGGTACAAA
2161 GACGAGAGGC TGCTGGAGGA AAAGTCTGGA GTCGACTTGG CGGACTCCAA CCAGAAGCTG
2221 AGCATCCAGC GCGTGCGCGA GGAGGATGCG GGACCGTATC TGTGCAGCGT GTGCAGACCC
2281 AAGGGCTGCG TCAACTCCTC CGCCAGCGTG GCCGTGGAAG GCTCCGAGGA TAAGGGCAGC
2341 ATGGAGATCG TGATCCTTGT CCGTACCGGC GTCATCGCTG TCTTCTTCTG GGTCTCTCTC
2401 CTCCTCATCT TCTGTAACAT GAGGAGGCCG GCCCACGCAG ACATCAAGAC GGGCTACCTG
2461 TCCATCATCA TGGACCCCGG GGAGGTGCCT CTGGAGGAGC AATGCGAATA CCTGTCTTAC
2521 GATGCCAGCC AGTGGGAATT CCCCCGAGAG CGGCTGCACC TGGGGAGAGT GCTCGGCTAC
2581 GCGCCTTTCG GGAAGGTGGT GGAAGCCTCC GCTTTCGGCA TCCACAAGGG CAGCAGCTGT
2641 GACACCGTGG CCGTGAAAAT GCTGAAAGAG GCGGCCACGG CCAGCGAGCA GCGCGCGCTG
2701 ATGTCGGAGC TCAAGATCCT CATTACATC GGCAACCACC TCAACGTGGT CAACCTCTC
2761 GGGGCGTGCA CCAAGCCGCA GGGCCCCCTC ATGGTGATCG TGGAGTTCTG CAAGTACGCG
2821 AACCTCTCCA ACTTCTTGGC CGCCAAGCGG GACGCCCTCA GCCCCTGCGC GGAGAAGTCT
2881 CCCGAGCAGC GCGGACGCTT CCGCGCCATG GTGGAGCTCG CCAGGCTGGA TCGGAGGCGG
2941 CCGGGGAGCA GCGACAGGGT CCTCTTCGCG CGGTTCTCGA AGACCGAGGG CGGAGCGAGG
3001 CGGGCTTCTC CAGACCAAGA AGCTGAGGAC CTGTGGCTGA GCCCCTGAC CATGGAAGAT
3061 CTTGTCTGCT ACAGCTTCCA GGTGGCCAGA GGGATGGAGT TCCTGGCTTC CCGAAAGTGC
3121 ATCCACAGAG ACCTGGCTGC TCGGAACATT CTGCTGTCGG AAAGCGACGT GGTGAAGATC
3181 TGTGACTTTG GCCTTGCCCG GGACATCTAC AAAGACCCCG ACTACGTCG CAAGGGCAGT

FIGURE 2A (PAGE 1 OF 2)

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3241 GCCCGGCTGC CCCTGAAGTG GATGGCCCCT GAAAGCATCT TCGACAAGGT GTACACCACG
3301 CAGAGTGACG TGTGGTCCTT TGGGGTGCTT CTCTGGGAGA TCTTCTCTCT GGGGGCCTCC
3361 CCGTACCCTG GGGTGCAGAT CAATGAGGAG TTCTGCCAGC GCGTGAGAGA CGGCACAAGG
3421 ATGAGGGCCC CGGAGCTGGC CACTCCCGCC ATACGCCACA TCATGCTGAA CTGCTGGTCC
3481 GGAGACCCCA AGGCGAGACC TGCATTCTCG GACCTGGTGG AGATCCTGGG GGACCTGCTC
3541 CAGGGCAGGG GCCTGCAAGA GGAAGAGGAG GTCTGCATGG CCCC GCGCAG CTCTCAGAGC
3601 TCAGAAGAGG GCAGCTTCTC GCAGGTGTCC ACCATGGCCC TACACATCGC CCAGGCTGAC
3661 GCTGAGGACA GCCCGCCAAG CCTGCAGCGC CACAGCCTGG CCGCCAGGTA TTACAACCTGG
3721 GTGTCCTTTC CCGGGTGCCT GGCCAGAGGG GCTGAGACCC GTGGTTCCTC CAGGATGAAG
3781 ACATTTGAGG AATTCCCCAT GACCCCAACG ACCTACAAAG GCTCTGTGGA CAACCAGACA
3841 GACAGTGGGA TGGTGCTGGC CTCGGAGGAG TTTGAGCAGA TAGAGAGCAG GCATAGACAA
3901 GAAAGCGGCT TCAGCTGTAA AGGACCTGGC CAGAATGTGG CTGTGACCAG GGCACACCCT
3961 GACTCCCAAG GGAGGCGGCG GCGGCCTGAG CGGGGGGCCG GAGGAGGCCA GGTGTTTTAC
4021 AACAGCGAGT ATGGGGAGCT GTCGGAGCCA AGCGAGGAGG ACCACTGCTC CCCGTCTGCC
4081 CGCGTGACTT TCTTCACAGA CAACAGCTAC TAA

10081126.022202

1 MQRGAALCLR LWLCLGLLDG LVSGYSMTTP TLNITEESHV IDTGDSLISIS CRGQHPLEWA
61 WPGAQEAPAT GDKDSEDGTV VRDCEGTAR PYCKVLLLHE VHANDTGSYV CYYKYIKARI
121 EGTTAASSYV FVRDFEQPFI NKPDTLLVNR KDAMWVPCLV SIPGLNVTLR SQSSVLWPDG
181 QEVVWDDRRG MLVSTPLLHD ALYLQCETTW GDQDFLSNPF LVHITGNELY DIQLLPRKSL
241 ELLVGEKLVL NCTVWAEFNS GVTFDWDYPG KQAERGWVP ERRSQQTHTE LSSILTIHNV
301 SQHDLGSYVC KANNGIQRF R ESTEVIVHEN PFISVEWLKG PILEATAGDE LVKLPVKLAA
361 YPPPEFQWYK DGKALSGRHS PHALVLKEVT EASTGTYTLA LWNSAAGLRR NISLELVVNV
421 PPQIHEKEAS SPSIYSRHSR QALTCTAYGV PLPLSIQWHW RPWTPCKMFA QRSLRRRQQQ
481 DLMPQCRDWR AVTTQDAVNP IESLDTWTEF VEGKNKTVSK LVIQNAVSA MYKCVVSNKV
541 GQDERLIYFY VTTIPDGFTI ESKPSEELLE GQPVLLSCQA DSYKYEHLRW YRLNLSTLHD
601 AHGNPLLLDC KNVHLFATPL AASLEEVAPG ARHATLSLSI PRVAPEHEGH YVCEVQDRRS
661 HDKXCHKKYL SVQALEAPRL TQNLTDLLVN VSDSLEMQL VAGAHAPSIV WYKDERLLEE
721 KSGVDLADSN QKLSIQRVRE EDAGRYLCSV CNAKGCVNSS ASVAVEGSED KGSMEIVILV
781 GTGVIAVFFW VLLLLIFCNM RRPADADIKT GYLSIIMDPG EVPLEEQCEY LSYDASQWEF
841 PRERLHLGRV LYGAFGKVV EASAFGIHKG SSCDTVAVKM LKEGATASEH RALMSELKIL
901 IHIGNHLNVV NLLGACTKPQ GPLMVIVEFC KYGNLSNFLR AKRDAFSPCA EKSPEQRGRF
961 RAMVELARLD RRRPGSSDRV LFRFSEKTEG GARRASPDQE AEDLWLSPLT MEDLVCYSFQ
1021 VARGMEFLAS RKCIHRDLAA RNILLSSESDV VKICDFGLAR DIYKDPDYVR KGSARLPLKW
1081 MAPESIFDKV YTTQSDVWSF GVLLWEIFSL GASPYPGVQI NEEFCQRLRD GTRMRAPELA
1141 TPAIRRIMLN CWSGDPKARP AFSELVEILG DLLQGRGLQE EEEVCMAPRS SQSSEEGSFS
1201 QVSTMALHIA QADAEDSPPS LQRHSLAARY YNWSFPGCL ARGAEETRGS RMKTFEEFPM
1261 TPTTYKGSVD NQTDSGMVLA SEEFQIESR HRQESGFSCK GPGQNVAVTR AHPDSQGRRR
1321 RPERGARGGQ VFYNSEYGEL SEPSEEDHCS PSARVTFFTD NSY

FIGURE 2B

20220 "Set" 1500T

10081.125 " 022202

1	CGCGGGGTGT	TCTGGTGTCC	CCCGCCCCGC	CTCTCCAAAA	AGCTACACCG	ACGCGGACCG
61	CGGCGGCGTC	CTCCCTCGCC	CTCGCTTCAC	CTCGCGGGCT	CCGAATGCGG	GGAGCTCGGA
121	TGTCCGGTTT	CCTGTGAGGC	TTTTACCTGA	CACCCGCCGC	CTTTCCCCGG	CACTGGCTGG
181	GAGGGCGCCC	TGCAAAGTTG	GGAACGCGGA	GCCCCGGACC	CGCTCCCGCC	GCCTCCGGCT
241	CGCCAGGGG	GGGTCGCCGG	GAGGAGCCCG	GGGGAGAGGG	ACCAGGAGGG	GCCCGCGGCC
301	TCGCAGGGGC	GCCCCGCGCC	CCACCCCTGC	CCCCGCCAGC	GGACCGGTCC	CCCACCCCCG
361	GTCCTTCCAC	CATGCACTTG	CTGGGCTTCT	TCTCTGTGGC	GTGTTCTCTG	CTCGCCGCTG
421	CGCTGCTCCC	GGGTCTCTCG	GAGGCGCCCG	CCGCCGCCGC	CGCCTTCGAG	TCCGGACTCG
481	ACCTCTCGGA	CGCGGAGCCC	GACGCGGGCG	AGGCCACGGC	TTATGCAAGC	AAAGATCTGG
541	AGGAGCAGTT	ACGGTCTGTG	TCCAGTGTAG	ATGAACTCAT	GACTGTACTC	TACCCAGAAT
601	ATTGGAAAA	TGACAAAGTGT	CAGCTAAGGA	AAGGAGGCTG	GCAACATAAC	AGAGAACAGG
661	CCAACCTCAA	CTCAAGGACA	GAAGAGACTA	TAAAATTTGC	TGCAGCACAT	TATAATACAG
721	AGATCTTGAA	AAGTATTGAT	AATGAGTGGA	GAAAGACTCA	ATGCATGCCA	CGGGAGGTGT
781	GTATAGATGT	GGGGAAGGAG	TTTGAGAGTCG	CGACAAACAC	CTTCTTTAAA	CCTCCATGTG
841	TGTCCGTCTA	CAGATGTGGG	GGTTGCTGCA	ATAGTGAGGG	GCTGCAGTGC	ATGAACACCA
901	GCACGAGCTA	CCTCAGCAAG	ACGTTATTTG	AAATTACAGT	GCCTCTCTCT	CAAGGCCCCA
961	AACCAGTAAC	AATCAGTTTT	GCCAATCACA	CTTCCTGCCG	ATGCATGTCT	AAACTGGATG
1021	TTTACAGACA	AGTTCATTCC	ATTATTAGAC	GTTCCCTGCC	AGCAACACTA	CCACAGTGTC
1081	AGGCAGCGAA	CAAGACCTGC	CCCACCAATT	ACATGTGGAA	TAATCACATC	TGCAGATGCC
1141	TGGCTCAGGA	AGATTTTATG	TTTTCTCTCG	ATGCTGGAGA	TGACTCAACA	GATGGATTCC
1201	ATGACATCTG	TGGACCAAAC	AAGGAGCTGG	ATGAAGAGAC	CTGTCAGTGT	GTCTGCAGAG
1261	CGGGGCTTCG	GCCTGCCAGC	TGTGACCCCC	ACAAAGAACT	AGACAGAAAC	TCATGCCAGT
1321	GTGTCTGTAA	AAACAAACTC	TTCCCCAGCC	AATGTGGGGC	CAACCGAGAA	TTTGATGAAA
1381	ACACATGCCA	GTGTGTATGT	AAAAGAACCT	GCCCCAGAAA	TCAACCCCTA	AATCCTGGAA
1441	AATGTGCCTG	TGAATGTACA	GAAAGTCCAC	AGAAATGCTT	GTAAAAGGA	AAGAAGTTCC
1501	ACCACCAAAC	ATGCAGCTGT	TACAGACGGC	CATGTACGAA	CCGCCAGAAG	GCTTGTGAGC
1561	CAGGATTTTC	ATATAGTGAA	GAAGTGTGTC	GTTGTGTCCC	TTCATATTGG	AAAAGACCAC
1621	AAATGAGCTA	AGATTGTACT	GTTTTCCAGT	TCATCGATTT	TCTATTATGG	AAAAGTGTGT
1681	TGCCACAGTA	GAAGTGTCTG	TGAACAGAGA	GACCCTTGTG	GGTCCATGCT	AACAAAGACA
1741	AAAGTCTGTC	TTTCCTGAAC	CATGTGGATA	ACTTTACAGA	AATGGACTGG	AGCTCATCTG
1801	CAAAAGGCCT	CTTGTAAGAA	CTGGTTTTCT	GCCAATGACC	AAACAGCCAA	GATTTTCCTC
1861	TTGTGATTTT	TTTAAAAGAA	TGACTATATA	ATTTATTTCC	ACTAAAAATA	TTGTTTCTGC
1921	ATTCATTTTT	ATAGCAACAA	CAATTGGTAA	AACTCACTGT	GATCAATATT	TTTATATCAT
1981	GCAAAATATG	TTTAAAATAA	AATGAAAATT	GTATT		

FIGURE 3A

MHLLGFFSVACSLLAALLPGPREAPAAAAAFESGLDLSDAEPDAGEATAYASKDLEEQLRSVSSVDELM
TVLYPEYWKMYKCQLRKGGWQHNREQANLNSRTEETIKFAAAHYNTEILKSIDNEWRTQCMPREVCIDV
GKEFGVATNTFFKPPCVSVYRCGGCCNSEGLQCMNTSTSYLSKTLFEITVPLSQGPKPVTISFANHTSCR
CMSKLDVYRQVHSIIRSLPATLPQCQAANKTCPTNYMWNHICRCLAQEDFMFSSDAGDDSTDGFHDIC
GPNKELDEETCQCVCRAGLRPASCGRPHKELDRNSCQCVCKNKLFPSCGANREFDENTCQCVCKRTCPRN
QPLNPGKCAECTESPQKCLLKGGKFHHQTCSCYRRPCTNRQKACEPGFSYSEVCRVPSYWKRPQMS

FIGURE 3B